

## SEQUENCE LISTING

<110> Martin Laforest  
 Nathalie Hubert  
 Benoit S. Landry

<120> Methods for Relative Quantification of  
 Specific Nucleic Acid Sequences

<130> 14187-1PCT

<160> 8

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 1600

<212> DNA

<213> E. coli

<400> 1

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tatccaatta gagtctcata ttcaactctca	atccaaataa tctgcaccgg atctggatcg	180
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<210> 2

<211> 1574

<212> DNA

<213> Brassica napus

<400> 2

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<212> DNA  
<213> Artificial Sequence

<220>  
<223> Sequence to be used as a primer

<400> 3  
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21

<210> 4  
<211> 20  
<212> DNA  
<213> Artificial Sequence

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<223> Sequence to be used as a primer

<400> 4  
gccatgggtc acgacgagat

20

<210> 5  
<211> 21  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> Sequence to be used as a primer

<400> 5  
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<210> 6  
<211> 21  
<212> DNA  
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<223> Sequence to be used as a primer

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<210> 7  
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<212> DNA  
<213> Artificial Sequence

<220>  
<223> Sequence to be used as a primer

<400> 7  
tttggaggag tgagtt 16

<210> 8  
<211> 16  
<212> DNA  
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<220>  
<223> Sequence to be used as a primer

<400> 8  
ctgccgagaa agtatac 16

<210> 9  
<211> 11  
<212> DNA  
<213> Artificial Sequence

<220>  
<223> This sequence is the dispension order of  
nucleotides given in Example 1 of the application.

<400> 9  
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